

# Exploring the plant microbiome for managing pathogens and resistances

**Gabriele Berg** Environmental Biotechnology TU Graz AUSTRIA



#### **INTRODUCTION:** Microbiome research – a new field in science





Biome: a reasonably well defined habitat which has distinct bio-physio-chemical properties

MICROBIOME = microbial community occupying a reasonably well defined habitat which has distinct physio-chemical properties. The term thus not only refers to the microorganisms involved but also encompasses their theatre of activity.

125 JM Whipps et al. 1988



#### **INTRODUCTION: Studying the fruit microbiome**



#### **PROBLEM: Biodiversity loss world-wide**



#### **Anthropocene:**

#### Human activity and intense agriculture caused:

- ✓ altered biogeochemical cycles
- ✓ species extinctions 100 to 1,000 times higher
- ✓ 1,000,000 species threatened with extinction



Science and Policy for People and Nature



## Increasing problems to control pathogens and resistances

#### **1.** Which microbial diversity is associated with plants (seed/fruit)?



#### 2. Is it possible to manage pathogens and resistances?

| Туре                            | Denomination           | Pedigree                                     | Geographic origin           | Seed samples |
|---------------------------------|------------------------|--|-----------------------------|--------------|
| Homozygous<br>inbred lines      | Line A - D             | -  | Austria                     | 默族           |
| Single cross<br>hybrid          | Gleisdorfer<br>Diamant | Line A x Line B                              | Austria                     | 528          |
| Three-way cross<br>hybrids      | GL Opal<br>GL Rustikal | Gl. Diamant x Line C<br>Gl. Diamant x Line D | Austria                     | e ko         |
| Population<br>cultivar          | GL Classic             |  | Austria                     |              |
| Single cross<br>zucchini hybrid | Naxos                  | -  | Netherlands                 | 350          |
| Segregating<br>breeding lines   | Line E – I             | -  | Germany,<br>Slovenia, China |              |

The microbiome was shaped by breeding The microbiome correlated with resistance against

Erwinia carotovora Enterobacteriaceae

[Adam et al. Plant and Soil 2016]









Bacteria on the petal of the withered female oil pumpkin flower visualized by CLSM and FISH. 3D construction by the software IMARIS.







Bacteria on the pollen grains on the pistil of the withered female oil pumpkin flower by CLSM and FISH.



### The Native Alpine Seed Microbiota

#### **1.** Which microbial diversity is associated with native plants?



#### **CONCLUSION:** The plant microbiome



#### **The Plant microbiome**

- changed during plants life cycle
- Is specific for plant species
- Was shaped by co-evolution
- Was shaped by breeding
- Has important function for the holo-biont
- Is vertically transmitted by seeds

### A healthy plant microbiome is highly diverse, rich and evenly structured.

[Berg & Raaijmakers ISME J 2018]



https://apfelmikrobiom.tugraz.at/kunst-konzept/



#### ✓ 10<sup>8</sup> 16S rRNA bacterial gene copy numbers were determined in each g apple

- ✓ fruit pulp and seeds were bacterial hot spots, while the peel was less colonized
- $\checkmark$  One apple a day provides more than 100 million bacteria independed of origin
- $\checkmark\,$  Organically produced apples contain a more diverse microbiome
- ✓ One apple a day provides more than 100 million bacteria



[Wassermann et al. Frontiers in Microbiology 2019]



| Ralstonia             | Sphingomonas   |
|-----------------------|--|
| Pseudomonas           | Massilia   |
| Methylobacterium      | Pantoea  |
| Burkholderia          | Rhizobiales sp.  |
| Hymenobacter          | Bacillus   |
| Variovorax            | Mucilaginibacter   |
| Curtobacterium        | Zymomonas  |
| Comamonadaceae sp.    | Spirosoma  |
| Acinetobacter         | Frondihabitans   |
| Pelomonas             | Acidiphilium   |
| Bdellovibrio          | Novosphingobium  |
| Flavobacterium        | Acetobacteraceae sp.   |
| Kineococcus           | Sphingomonadaceae sp.  |
| Planctomycetes sp.    | Erwinia  |
| Myxococcales sp.      | Rhizobium  |
| Armatimonadetes sp.   | Amnibacterium  |
| Staphylococcus        | Deinococcus  |
| Oligoflexales sp.     | Caenimonas   |
| Oxalobacteraceae sp.  | Sphingobium  |
| Pedobacter            | Sorangium  |
| Rathayibacter         | Gluconobacter  |
| Legionella            | Aquabacterium  |
| Bradyrhizobium        | Corynebacterium  |
| Paenibacillus         | Acidobacteria sp.  |
| Microbacteriaceae sp. | Chryseobacterium   |
| Rhizobiales sp.       | Terriglobus  |
| Pectobacterium        | Stenotrophomonas   |
| Nitrospira            | Soil Crenarchaeotic Group(SCG)   |
| III Buchnera          | Arthrobacter   |
| Streptococcus         | Reyranella   |
| Fructobacillus        | and a second state of the second state of the second second second second second second second second second s |



[Wassermann *et al.* Frontiers in Microbiology 2019]



[Wassermann et al. Frontiers in Microbiology 2019]

#### **Sparkling science: translating science**















#### Microbial diversity on apples is of international interest



Apter sand nicht nur gesund, wen sie vitamine enthalten. Dies diest beherbergt auch eine grobe Anzahl Mikzoorganismen, die nach dem Verzehr den Darm besiedeln. Biofrüchte enthalten andere Bakterien als konventionell bergestellte.

#### **PROBLEM:** Food loss

#### **Global food loss**



#### 2. Is it possible to manage microbial biodiversity?

#### 2. Is it possible to manage microbial biodiversity?



#### **2. Is it possible to manage microbial biodiversity?**





Gesellschaft für Biotechnologie mbH



#### **2. Is it possible to manage microbial biodiversity?**



[unpublished results]

#### 2. Is it possible to manage antimicrobial resistances?



A 50% reduction in diversity on abiotic surfaces led to a 20% increase in antibiotic resistances (P=0.01).

Increased confinement and cleaning resulted in:

- a loss of microbial diversity by maintained abundance
- a shift at superkindom level and from Gram-positive to Gram-negatives
- a shift within pan-genomes (Acinetobacter)
- a larger number of resistance genes

#### **CONCLUSION: Biodiversity loss world-wide**



Plant microbial diversity is a key for all health issues.

The microbiome connects our world.

The microbiome is important for one health issues



#### The plant microbiota is crucial for one health issues

